

SEQUENCE LISTING

TEOP CENTER TOO SOO

- <110> Hauptmann, Rudolph Himmler, Adolph Maurer-Fogy, Ingrid Stratowa, Christian
- <120> TNF Receptors, TNF Binding Proteins and DNAs Coding for Them
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- <140> 09/525,998
- <141> 2000-03-15
- <:150> 08/383,676
- <151> 1995-02-01
- <150> 08/153,287
- <151> 1993-11-17
- <150> 07/821,750
- <151> 1993-01-02
- <150> 07/511,430
- <151> 1990-04-20
- <160> 97
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aag aac cag tac Lys Asn Gln Tyr 130	Arg His '				
aat tgc agc ctc Asn Cys Ser Leu 145					
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Lys Asn Gln 130	Tyr Arg		yr Trp 35	Ser Glu		Leu F 140	Phe (Gln	Cys	Phe	
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Lys Gln Asn	Thr Val 165	Cys T	hr Cys	His Ala		Phe I	Phe	Leu	Arg 175	Glu	
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						cag Gln										336
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-						tgc Cys										519
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Met 1 Pro)> 10 Leu Gln	Val Gly	Pro Lys 20	His 5 Tyr	Leu Ile		Pro	Gln 25	10 Asn	Asn	Ser	Ile	Cys 30	15 Cys	Thr	
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Met 1 Pro Lys Gln Glu 65	Cys Asp 50 Asn	Val Gly His 35 Thr	Pro Lys 20 Lys Asp	His 5 Tyr Gly Cys	Leu Ile Thr Arg	His Tyr Glu 55	Pro Leu 40 Cys	Gln 25 Tyr Glu Ser	10 Asn Asn Ser Cys	Asn Asp Gly Ser 75	Ser Cys Ser 60 Lys	Ile Pro 45 Phe Cys	Cys 30 Gly Thr	15 Cys Pro Ala Lys	Thr Gly Ser Glu 80	
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Met 1 Pro Lys Gln 65 Met Cys	Cys Asp 50 Asn Gly	Val Gly His 35 Thr Gln Cys	Pro Lys 20 Lys Asp Leu Val Arg	His 5 Tyr Gly Cys Arg Glu 85 Lys	Leu Ile Thr Arg His 70 Ile Asn	His Tyr Glu 55 Cys	Pro Leu 40 Cys Leu Ser	Gln 25 Tyr Glu Ser Cys Arg 105	10 Asn Asn Ser Cys Thr 90 His	Asn Asp Gly Ser 75 Val	Ser Cys Ser 60 Lys Asp	Ile Pro 45 Phe Cys Arg	Cys 30 Gly Thr Arg Asp	15 Cys Pro Ala Lys Thr 95 Asn	Thr Gly Ser Glu 80 Val	

Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser 145 150 155 160
Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn 165 170
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tgt ccc caa gga aaa tat atc cac cct caa aat aat tcg att tgc tgt Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys 35 40 45
acc aag tgc cac aaa gga acc tac ttg tac aat gac tgt cca ggc ccg Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro 50 50 60
ggg cag gat acg gac tgc agg gag tgt gag agc ggc tcc ttc acc gct Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala 65 70 75 80
tca gaa aac cac ctc aga cac tgc ctc agc tgc tcc aaa tgc cga aag 288 Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys 85 90 95
gaa atg ggt cag gtg gag atc tct tct tgc aca gtg gac cgg gac acc Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr 100 105 110
gtg tgt ggc tgc agg aag aac cag tac cgg cat tat tgg agt gaa aac Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn 115 120 125

ctt ttc cag tgc ttc aat tgc agc ctc tgc ctc aat ggg acc gtg cac Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His

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Cys	Pro	Gln 35	Gly	Lys	Tyr	Ile	His 40	Pro	Gln	Asn	Asn	Ser 45	Ile	Cys	Cys	
Thr	Lys 50	Cys	His	Lys	Gly	Thr 55	Tyr	Leu	Tyr	Asn	Asp 60	Cys	Pro	Gly	Pro	
Gly 65	Gln	Asp	Thr	Asp	Cys 70	Arg	Glu	Cys	Glu	Ser 75	Gly	Ser	Phe	Thr	Ala 80	
Ser	Glu	Asn	His	Leu 85	Arg	His	Cys	Leu	Ser 90	Cys	Ser	Lys	Cys	Arg 95	Lys	
Glu	Met	Gly	Gln 100	Val	Glu	Ile	Ser	Ser 105	Cys	Thr	Val	Asp	Arg 110	Asp	Thr	
Val	Cys	Gly 115	Cys	Arg	Lys	Asn	Gln 120	Tyr	Arg	His	Tyr	Trp 125	Ser	Glu	Asn	
Leu	Phe 130	Gln	Cys	Phe	Asn	Cys 135	Ser	Leu	Cys	Leu	Asn 140	Gly	Thr	Val	His	
Leu 145	Ser	Cys	Gln	Glu	Lys 150	Gln	Asn	Thr	Val	Cys 155	Thr	Cys	His	Ala	Gly 160	
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Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn

180

185

Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile

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<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: recombinant
 TNF-BP sequence

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Ser Ile Cys Cys Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp 20 25 30

Cy's Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly
35 40 45

Ser Phe Thr Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser 50 55 60

Lys Cys Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val
65 70 75 80

Asp Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr 85 90 95

Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn 100 105 110

Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr 115 120 125

Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser 130 135 140

Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile 145 150 155 160

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: cDNA insert of lambdaTNF-BP15 and pTNF-BP15 vectors

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aatg	igācē	gag t	gaga	aggco	ca ta	agcto	gtato	g gc						gtg Val		233
														ata Ile		281
		-												gag Glu		329
														aat Asn		377
														aat Asn 70		425
														agc Ser		473
			_		_									tgc Cys		521
aaa Lys	tgc Cys 105	cga Arg	aag Lys	gaa Glu	atc Ile	ggt Gly 110	cag Gln	gtg Val	gag Glu	atc Ile	tct Ser 115	tct Ser	tgc Cys	aca Thr	gtg Val	569
														cat His		617
														ctc Leu 150		665
														tgc Cys		713

						cta Leu										761
						gag Glu 190										809
						gag Glu										857
						ctt Leu										905
						cgg Arg										953
tgt Cys	ggg Gly	aaa Lys 250	tcg Ser	aca Thr	cct Pro	gaa Glu	aaa Lys 255	gag Glu	ggg Gly	gag Glu	ctt Leu	gaa Glu 260	gga Gly	act Thr	act Thr	1001
						aac Asn 270										1049
						agt Ser										1097
						ggt Gly										1145
						tat Tyr										1193
gcc Ala	ctc Leu	gcc Ala 330	tcc Ser	gac Asp	ccc Pro	atc Ile	ccc Pro 335	aac Asn	ccc Pro	ctt Leu	cag Gln	aag Lys 340	tgg Trp	gag Glu	gac Asp	1241
agc Ser	gcc Ala 345	cac His	aag Lys	cca Pro	cag Gln	agc Ser 350	cta Leu	gac Asp	act Thr	gat Asp	gac Asp 355	ccc Pro	gcg Ala	acg Thr	ctg Leu	1289
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<212> PRT

<213> Artificial Sequence

<220>

<400> 22

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Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly Leu Val Pro
20 25 30

His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro Gln Gly Lys 35 40 45

Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys 50 55 60

Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln Asp Thr Asp 65 70 75 80

Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu Asn His Leu 85 90 95

Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Ile Gly Gln Val

Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys Gly Cys Arg 115 120 125

Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe 130 135 140

Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln Glu 145 150 155 160

Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu 165 170 175

Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Arg 180 185 190

Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp Ser 195 200 205

Gly Thr Thr Val Leu Leu Pro Leu Val Ile Phe Phe Gly Leu Cys Leu 210 215 220

Leu Ser Leu Leu Phe Ile Gly Leu Met Tyr Arg Tyr Gln Arg Trp Lys 235 230 235

Ser Lys Leu Tyr Ser Ile Val Cys Gly Lys Ser Thr Pro Glu Lys Glu 245 250 255

Gly Glu Leu Glu Gly Thr Thr Thr Lys Pro Leu Ala Pro Asn Pro Ser 260 265 270

Phe Ser Pro Thr Pro Gly Phe Thr Pro Thr Leu Gly Phe Ser Pro Val 280 Pro Ser Ser Thr Phe Thr Ser Ser Ser Thr Tyr Thr Pro Gly Asp Cys 295 Pro Asn Phe Ala Ala Pro Arg Arg Glu Val Ala Pro Pro Tyr Gln Gly 310 305 Ala Asp Pro Ile Leu Ala Thr Ala Leu Ala Ser Asp Pro Ile Pro Asn 330 Pro Leu Gln Lys Trp Glu Asp Ser Ala His Lys Pro Gln Ser Leu Asp 340 345 Thr Asp Asp Pro Ala Thr Leu Tyr Ala Val Val Glu Asn Val Pro Pro 360 Leu Arg Trp 370 <210> 23 <211> 6414 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: pADCMV1 vector <220> <221> unsure <222> (344) <223> "n" can be a, g, c, or t <220> <221> unsure <222> (4157) <223> "n" can be a, g, c, or t <220> <221> unsure <222> (5135) <223> "n" can be a, g, c, or t <220> <221> unsure <222> (6255) <223> "n" can be a, g, c, or t tcgacattga ttattgacta gttattaata gtaatcaatt acggggtcat tagttcatag 60 cccatatatq qaqttccqcq ttacataact tacggtaaat ggcccgcctc gctgaccgcc 120

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ctg gct ctg ctg Leu Ala Leu Leu						337
cct tct ctt ggt Pro Ser Leu Gly 35						385
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gtg gag att tct Val Glu Ile Ser 115						625
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gtg gac tgc agc Val Asp Cys Ser 145		Asn Gly	Thr Val T			721
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tca ggt act gcc Ser Gly Thr Ala						913

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								gta Val	1201
								tcc Ser	1249
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								ctg Leu	1345
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								cag Gln	1441
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								cgc Arg 430	1537
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Tyr Ala His Pro Lys Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys
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Gly Thr Tyr Leu Val Ser Asp Cys Pro Ser Pro Gly Gln Glu Thr Val 65 70 75 80

Cys Glu Leu Ser His Lys Gly Thr Phe Thr Ala Ser Gln Asn His Val 85 90 95

Arg Gln Cys Leu Ser Cys Lys Thr Cys Arg Lys Glu Met Phe Gln Val

Glu Ile Ser Pro Cys Lys Ala Asp Met Asp Thr Val Cys Gly Cys Lys

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Lys	Leu	Cys 195	Leu	Pro	Pro	Val	Ala 200	Asn	Val	Thr	Asn	Pro 205	Gln	Asp	Ser
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521

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Lys	Leu	Cys 195	Leu	Pro	Gln	Ile	Glu 200	Asn	Val	Lys	Gly	Thr 205	Glu	Asp	Ser
Gly	Thr 210	Thr	Val	Leu	Leu	Pro 215	Leu	Val	Ile	Phe	Phe 220	Gly	Leu	Cys	Leu
Leu 225	Ser	Leu	Leu	Phe	Ile 230	Gly	Leu	Met	Tyr	Arg 235	Tyr	Gln	Arg	Trp	Lys 240
Ser	Lys	Leu	Tyr	Ser 245	Ile	Val	Cys	Gly	Lys 250	Ser	Thr	Pro	Glu	Lys 255	Glu
Gly	Glu	Leu	Glu 260	Gly	Thr	Thr	Thr	Lys 265	Pro	Leu	Ala	Pro	Asn 270	Pro	Ser

Phe Ser Pro Thr Pro Gly Phe Thr Pro Thr Leu Gly Phe Ser Pro Val

275 280 285

Pro Ser Ser Thr Phe Thr Ser Ser Ser Thr Tyr Thr Pro Gly Asp Cys 290 295 300

Pro Asn Phe Ala Ala Pro Arg Arg Glu Val Ala Pro Pro Tyr Gln Gly 305 310 315 320

Ala Asp Pro Ile Leu Ala Thr Ala Leu Ala Ser Asp Pro Ile Pro Asn 325 330 335

Pro Leu Gln Lys Trp Glu Asp Ser Ala His Lys Pro Gln Ser Leu Asp 340 345 350

Thr Asp Asp Pro Ala Thr Leu Tyr Ala Val Val Glu Asn Val Pro Pro 355 360 365

Leu Arg Trp Lys Glu Phe Val Arg Arg Leu Gly Leu Ser Asp His Glu 370 375 380

Ile Asp Arg Leu Glu Leu Gln Asn Gly Arg Cys Leu Arg Glu Ala Gln 385 390 395 400

Tyr Ser Met Leu Ala Thr Trp Arg Arg Arg Thr Pro Arg Arg Glu Ala 405 410 415

Thr Leu Glu Leu Leu Gly Arg Val Leu Arg Asp Met Asp Leu Leu Gly 420 425 430

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cleavage peptide

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gat acg gac tgc agg gag tgt gag agc ggc tcc ttc aca gcc tca gaa Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu 35 40 45	144
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Asn Asn Lys
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EBI-2467

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